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Preliminary Amendment
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CLAIMS:

1-65 (canceled)

- 66. (new) An amino acid molecule comprising a peptide comprising at least one of the following characteristics:
 - (a) being capable binding to ManLAM binding antibodies; and
 - (b) being capable of eliciting, upon immunization in a subject, production of ManLAM-binding antibodies.
- 67. (new) The amino acid molecule of Claim 66, wherein said ManLAM binding antibodies are anti-ManLAM antibodies.
- 68. (new)The amino acid molecule of Claim 66, wherein said ManLAM binding antibodies are monoclonal antibodies (mAbs).
- 69. (new) The amino acid molecule of Claim 68, wherein said mAbs are CS40 antibodies.
- 70. (new) The amino acid molecule of Claim 66, which does not bind to antibodies directed against lipoglycans selected from non-mannolsylated and low mannosylated lipoglycans.
- 71. (new) The amino acid molecule of Claim 70, which does not bind to CS35 anti-LAM mAb, 735 anti-ploy $\alpha(2\rightarrow 8)$ N-acetyl neuraminic acid mAb, and 2H1 anti- glucuronoxylomannan mAb.
- 72. (new) The amino acid molecule of Claim 66, wherein said peptide has an internal aromatic amino acid residue.
- 73. (new) The amino acid molecule of Claim 72, wherein said aromatic amino acid residue is selected from Phenylalanine (F), Histidine (H), Tryptophan (W), Tyrosine (Y) and conservative substitutions thereof.
- 74. (new) The amino acid molecule of Claim 72, wherein said peptide comprises a hydrophilic amino acid residue adjacent and preceding said aromatic residue.

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75. (new) The amino acid molecule of Claim 72, wherein said aromatic residue is W.

76. (new) The amino acid molecule of Claim 66, wherein said peptide comprises the sequence selected from:

ISLTEWSMWYRH (SEQ ID NO:1);
EEGPWSTHVGRT (SEQ ID NO:2);
WGNEGGDHLQPV (SEQ ID NO:3);

SLKIRWELKMYQE (SEQ ID NO:4);

AVERWEKHTWSE (SEQ ID NO:5);

and immunologic modifications thereof.

- 77. (new) The amino acid molecule of Claim 66, wherein said peptide comprises the sequence ISLTEWSMWYRH (SEQ ID NO:1), or an immunogenic modification thereof.
- 78. (new) A method for diagnosing a mycobacterial infection in a subject the method comprising:
 - (a) contacting said sample with an amino acid molecule comprising a peptide comprising at least one of the following characteristics:
 - i) being capable of binding to ManLAM-binding antibodies, and
 - ii) being capable of eliciting, upon immunization in a subject, production of ManLAM binding antibodies; and
 - (b) determining formation of a complex comprising said amino acid molecule and ManLAM binding antibodies, if present in the sample,

wherein a positive determination indicates mycobacterial infection in the subject.

79. (new) A method for determining whether a subject has active mycobacterial infection the method comprising:

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- (a) contacting a sample from said subject with an amino acid molecule comprising a peptide comprising at least one of the following characteristics:
 - i) being capable of binding to ManLAM-binding antibodies, and
 - ii) being capable of eliciting, upon immunization in a subject, production of ManLAM binding antibodies;
- (b) determining level of complexes comprising said amino acid molecule and ManLAM binding antibodies; and
- (c) comparing said level to a standard, wherein a level higher than the standard indicates active myobacterial infection in the subject.
- 80. (new) A method for determining treatment efficacy in a subject having a mycobacterial infection, the method comprising:
 - (a) contacting samples from said subject, from at least two discrete time points, with an amino acid molecule comprising a peptide comprising at least one of the following characteristics:
 - i) being capable of binding to ManLAM-binding antibodies, and
 - ii) being capable of eliciting, upon immunization in a subject, production of ManLAM binding antibodies; and
 - (b) determining level of complexes comprising said amino acid molecule and ManLAM binding antibodies in said samples,

wherein a difference in the level between the two time points is indicative of the effectiveness of the treatment.

- 81. (new) A kit for diagnosing mycobacterial infection in a subject comprising an amino acid molecule comprising a peptide, the peptide comprising at least one of the following characteristics:
 - (a) being capable of binding to ManLAM-binding antibodies; and
 - (b) being capable of eliciting production of ManLAM binding antibodies.

- 82. (new) A vaccine comprising an immunologically acceptable carrier and as an active agent an amino acid molecule comprising a peptide comprising at least one of the following characteristics:
 - (a) being capable of binding to ManLAM-binding antibodies; and
 - (b) being capable of eliciting, upon immunization of a subject, production of ManLAM binding antibodies.
- 83. (new) The vaccine of Claim 82, wherein said ManLAM binding antibodies are anti-ManLAM antibodies.
- 84. (new) The vaccine of Claim 83, wherein the amino acid molecule does not bind to antibodies directed against lipoglycans selected from non-mannolsylated and low mannosylated lipoglycans.
- 85. (new) The vaccine of Claim 84, which amino acid molecule does not bind to CS35 anti-LAM mAb, 735 anti-ploy $\alpha(2\rightarrow 8)$ N-acetyl neuraminic acid mAb, and 2H1 anti- glucuronoxylomannan mAb.
- 86. (new) The vaccine of Claim 83, wherein said amino acid molecule comprises at least one peptide having an internal aromatic amino acid residue.
- 87. (new) The vaccine of Claim 86, wherein said aromatic amino acid residue is selected from Phenylalanine (F), Histidine (H), Tryptophan (W), Tyrosine (Y) and conservative substitutions thereof.
- 88. (new) The vaccine of Claim 86, wherein said peptide comprises a hydrophilic amino acid residue adjacent and preceding said aromatic residue.
- 89. (new) The vaccine of Claim 88, wherein said aromatic residue is W.
- 90. (new) The vaccine of Claim 83, wherein said amino acid molecule comprises at least one peptide having the sequence selected from:

ISLTEWSMWYRH (SEQ ID NO:1);

EEGPWSTHVGRT (SEQ ID NO:2);

WGNEGGDHLQPV (SEQ ID NO:3);

SLKIRWELKMYQE (SEQ ID NO:4);

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AVERWEKHTWSE (SEQ ID NO:5);

and immunologic modifications thereof.

- 91. (new) The vaccine of Claim 83, wherein amino acid comprises a peptide having the sequence ISLTEWSMWYRH (SEQ ID NO:1), and immunologic modifications thereof.
- 92. (new) A method of immunization of a subject against mycobacterial infection, the method comprises providing said subject with an immunizing amount of an amino acid molecule comprising a peptide comprising at least one of the following characteristics:
 - (a) being capable of binding to ManLAM-binding antibodies; and
 - (b) being capable of eliciting, upon immunization of a subject, production of ManLAM binding antibodies.
- 93. (new) The method of Claim 92, wherein the amino acid molecule does not bind to antibodies directed against lipoglycans selected from non-mannolsylated and low mannosylated lipoglycans.
- 94. (new) The method of Claim 93, wherein the amino acid molecule does not bind to CS35 anti-LAM mAb, 735 anti-ploy $\alpha(2\rightarrow 8)$ N-acetyl neuraminic acid mAb, and 2H1 anti- glucuronoxylomannan mAb.
- 95. (new) The method of Claim 94, wherein said amino acid molecule comprises at least one peptide having an internal aromatic amino acid residue.
- 96. (new) The method of Claim 95, wherein said aromatic amino acid residue is selected from Phenylalanine (F), Histidine (H), Tryptophan (W), Tyrosine (Y) and conservative substitutions thereof.
- 97. (new) The method of Claim 95, wherein said peptide comprises a hydrophilic amino acid residue adjacent and preceding said aromatic residue.
- 98. (new) The method of Claim 95, wherein said aromatic residue is W.
- 99. (new) The method of Claim 93, wherein said amino acid molecule comprises at least one peptide having the sequence selected from:

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ISLTEWSMWYRH (SEQ ID NO:1);

EEGPWSTHVGRT (SEQ ID NO:2);

WGNEGGDHLQPV (SEQ ID NO:3);

SLKIRWELKMYQE (SEQ ID NO:4);

AVERWEKHTWSE (SEQ ID NO:5);

and immunologic modifications thereof.

100. (new) The method of Claim 99, wherein the amino acid molecule comprises a peptide having the sequence ISLTEWSMWYRH (SEQ ID NO:1), and immunologic modification thereof.